

RAW SEQUENCE LISTING FMARK OF ERROR REPORT



The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: /0/660, 893
Source: /F49
Date Processed by STIC: ///04

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO-REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER

VERSION 4.2 PROGRAM, ACCESSIBLE THROUGHTHE: U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker/chkrnote.htm

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: Commissioner for Patents, P.O. Box-1450, Alexandria, VA 22313-1450

Revised 05/17/04

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JUL 1 6 2004

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Original die date
respend.
BA 7/21/04



1 2 2004	Raw Sequence Listing Error Summary
ERROR DETECTED	SUGGESTED CORRECTION SERIAL NUMBER: 10/660, 893
DEMADO ATTN: NEW RULES CASES	: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE
IWrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."
2Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.
Misaligned Amino Numbering	The numbering under each 5th amino acid is misaligned. Do not use tableodes between numbers: use space characters, instead, an interest of the space characters, instead, and interest of the space characters.
4Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.
5Variable Length	Sequence(s) contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.
6PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.
7Skipped Sequences (OLD RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped
	Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.
8Skipped Sequences (NEW RULES)	Sequence(s) missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000
9Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.
10_V Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence
Use of <220>	Sequence(s) missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 06/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)
Patentin 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.
3 Misuse of n/Xaa	"n" can only represent a single nucleotide; "Xaa" can only represent a single amino acid

AMC - Biotechnology Systems Branch - 09/09/2003





IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/660,893

DATE: 07/01/2004

TIME: 13:09:23

Input Set : A:\New Link, 0003.ST25.txt
Output Set: N:\CRF4\07012004\J660893.raw

```
3 <110> APPLICANT: Link, Charles
 5 <120> TITLE OF INVENTION: Methods and Compositions for Elucidating Protein Expression
         Profiles in Cells
 6
 8 <130> FILE REFERENCE: 05237.0003.CPUS00
10 <140> CURRENT APPLICATION NUMBER: US 10/660,893
11 <141> CURRENT FILING DATE: 2003-09-12
13 <160> NUMBER OF SEQ ID NOS: 5
15 <170> SOFTWARE: PatentIn version 3.2
                         invalid (2137 response. Sel iten 10 on Evan Summary
Does Not Comply Sheet.
17 <210> SEO ID NO: 1
18 <211> LENGTH: 9
19 <212> TYPE: PRT
20 <213 > ORGANISM: Synthetic
                                                         Corrected Diskette Needec
22 <400> SEQUENCE:
24 Tyr Pro Tyr Asp Val Pro Asp Tyr Ala
25 1
28 <210> SEQ ID NO: 2
29 <211> LENGTH: 10
30 <212> TYPE: PRT/
31 <213> ORGANISM: Synthetic
33 <400> SEQUENCE: 2
35 Glu Gln Lys Leu Ile Ser Glu Glu Asp Leu
39 <210> SEQ ID NO: 3
40 <211> LENGTH: 8
41 <212> TYPE: PRT
42 <213> ORGANISM: Synthetic
44 <400> SEQUENCE: 3
46 Asp Tyr Lys Asp Asp Asp Lys
47 1
50 <210> SEQ ID NO: 4
51 <211> LENGTH: 375
52 <212> TYPE: DNA
53 <213> ORGANISM: Synthetic
56 <220> FEATURE:
57 <221> NAME/KEY: misc_feature
58 <222> LOCATION: (3)..(4)
59 <223> OTHER INFORMATION: n is a, c, g, or t
61 <220> FEATURE:
62 <221> NAME/KEY: misc feature
63 <222> LOCATION: (8)..(8)
                                                      REC'D HOWREY SIMON ARNOLD & WHITE
64 <223> OTHER INFORMATION: n is a, c, g, or t
66 <220> FEATURE:
67 <221> NAME/KEY: misc_feature
```

JUL 1 6 2004

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RAW SEQUENCE LISTING DATE: 07/01/2004 PATENT APPLICATION: US/10/660,893 TIME: 13:09:23 Input Set: A:\New Link, 0003.ST25.txt Output Set: N:\CRF4\07012004\J660893.raw

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68 <222> LOCATION: (350)..(350)
  69 <223> OTHER INFORMATION: n is a, c, g, or t
  71 <220> FEATURE:
  72 <221> NAME/KEY: misc_feature
  73 <222> LOCATION: (358)..(374)
  74 <223> OTHER INFORMATION: n is a, c, g, or t
  76 <400> SEQUENCE: 4
-> 77 ttnnccgnga aageteeteg ceettgetea ceatgggatg ceattteeta ggtetgeete
  79 ttggccgttt ttctccaatg gtctctgctt tcttctgggc tgctttagag gggctcttgt
  81 tittgetgee tittgggtett cetetgggte tettaggaga gggeteacag gitggetett
  83 getgetgett cetgggtegg cegegteete gettetgtgg caceggggeg geaggttgte.
  85 cetgggetga tgtggaegge tgeeeggege ceteacegeg tgegeteate etgeeteeeg
-> 87 ccgccgctac cactgcctct cttttttttt ttttttttt ttttttgaaan ccccgggnnn
-> 89 nnnnnnnnn nnnc
  92 <210> SEQ ID NO: 5
 93 <211> LENGTH: 333
  94 <212> TYPE: DNA
  95 <213> ORGANISM: (Synthetic
  98 <220> FEATURE:
  99 <221> NAME/KEY: misc_feature
  100 <222> LOCATION: (3)..(3)
  101 <223> OTHER INFORMATION: n is a, c, g, or t
  103 <220> FEATURE:
  104 <221> NAME/KEY: misc_feature
  105 <222> LOCATION: (11) .. (11)
  106 <223> OTHER INFORMATION: n is a, c, g, or t
  108 <220> FEATURE:
  109 <221> NAME/KEY: misc_feature
  110 <222> LOCATION: (106)..(106)
  111 <223> OTHER INFORMATION: n is a, c, g, or t
  113 <220> FEATURE:
  114 <221> NAME/KEY: misc feature
  115 <222> LOCATION: (116)..(116)
  116 <223> OTHER INFORMATION: n is a, c, g, or t
  118 <220> FEATURE:
  119 <221> NAME/KEY: misc feature
  120 <222> LOCATION: (168)..(168)
  121 <223> OTHER INFORMATION: n is a, c, g, or t
  123 <220> FEATURE:
  124 <221> NAME/KEY: misc feature
  125 <222> LOCATION: (179)..(179)
  126 <223> OTHER INFORMATION: n is a, c, g, or t
  128 <220> FEATURE:
  129 <221> NAME/KEY: misc feature
  130 <222> LOCATION: (204)..(204)
  131 <223> OTHER INFORMATION: n is a, c, g, or t
  133 <220> FEATURE:
  134 <221> NAME/KEY: misc feature
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135 <222> LOCATION: (221)..(221)

RAW SEQUENCE LISTING DATE: 07/01/2004 PATENT APPLICATION: US/10/660,893 TIME: 13:09:23

Input Set : A:\New Link, 0003.ST25.txt
Output Set: N:\CRF4\07012004\J660893.raw

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136 <223 > OTHER INFORMATION: n is a, c, q, or t
138 <220> FEATURE:
139 <221> NAME/KEY: misc feature
140 <222> LOCATION: (224)..(224)
141 <223> OTHER INFORMATION: n is a, c, g, or t
143 <220> FEATURE:
144 <221> NAME/KEY: misc_feature
145 <222> LOCATION: (231)..(231)
146 <223> OTHER INFORMATION: n is a, c, g, or t
148 <220> FEATURE:
149 <221> NAME/KEY: misc feature
150 <222> LOCATION: (254)..(254)
151 <223> OTHER INFORMATION: n is a, c, g, or t
153 <220> FEATURE:
154 <221> NAME/KEY: misc_feature
155 <222> LOCATION: (272)..(272)
156 <223> OTHER INFORMATION: n is a, c, g, or t
158 <220> FEATURE:
159 <221> NAME/KEY: misc feature
160 <222> LOCATION: (275)..(275)
161 <223> OTHER INFORMATION: n is a, c, g, or t
163 <220> FEATURE:
164 <221> NAME/KEY: misc_feature
165 <222> LOCATION: (282)..(282)
166 <223> OTHER INFORMATION: n is a, c, g, or t
168 <220> FEATURE:
169 <221> NAME/KEY: misc_feature
170 <222> LOCATION: (285)..(286)
171 <223> OTHER INFORMATION: n is a, c, g, or t
173 <220> FEATURE:
174 <221> NAME/KEY: misc_feature
175 <222> LOCATION: (289)..(289)
176 <223> OTHER INFORMATION: n is a, c, g, or t
178 <220> FEATURE:
179 <221> NAME/KEY: misc_feature
180 <222> LOCATION: (292)..(293)
181 <223> OTHER INFORMATION: n is a, c, g, or t
183 <220> FEATURE:
184 <221> NAME/KEY: misc_feature
185 <222> LOCATION: (296)..(296)
186 <223> OTHER INFORMATION: n is a, c, g, or t
188 <220> FEATURE:
189 <221> NAME/KEY: misc_feature
190 <222> LOCATION: (299)..(299)
191 <223> OTHER INFORMATION: n is a, c, g, or t
193 <220> FEATURE:
194 <221> NAME/KEY: misc_feature
195 <222> LOCATION: (301)..(301)
196 <223> OTHER INFORMATION: n is a, c, g, or t
```

RAW SEQUENCE LISTING

DATE: 07/01/2004

PATENT APPLICATION: US/10/660.893

TIME: 13:09:23

Input Set : A:\New Link, 0003.ST25.txt Output Set: N:\CRF4\07012004\J660893.raw

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198 <220> FEATURE:
 199 <221> NAME/KEY: misc_feature
 200 <222> LOCATION: (304)..(304)
 201 <223> OTHER INFORMATION: n is a, c, g, or t
 203 <220> FEATURE:
 204 <221> NAME/KEY: misc feature
 205 <222> LOCATION: (306)..(308)
 206 <223> OTHER INFORMATION: n is a, c, g, or t
 208 <220> FEATURE:
 209 <221> NAME/KEY: misc feature
 210 <222> LOCATION: (311)..(311)
 211 <223> OTHER INFORMATION: n is a, c, g, or t
 213 <220> FEATURE:
 214 <221> NAME/KEY: misc feature
 215 <222> LOCATION: (326)..(327)
 216 <223> OTHER INFORMATION: n is a, c, g, or t
 218 <220> FEATURE:
 219 <221> NAME/KEY: misc_feature
 220 <222> LOCATION: (329)..(329)
 221 <223> OTHER INFORMATION: n is a, c, g, or t
 223 <220> FEATURE:
 224 <221> NAME/KEY: misc_feature
 225 <222> LOCATION: (331)..(331)
 226 <223> OTHER INFORMATION: n is a, c, g, or t
 228 <220> FEATURE:
 229 <221> NAME/KEY: misc_feature
 230 <222> LOCATION: (333)..(333)
 231 <223> OTHER INFORMATION: n is a, c, g, or t
 233 <400> SEQUENCE: 5
> 234 tengegacea netectegee ettgeteace atgggatget eeeggtggtg ggteggtggt
> 236 ccctgggcag gggtctccaa atcccggacg agcccccaaa tgaaanaccc ccgtcntggg
> 238 tagtcaatca ctcagaggag acceteceaa ggaacagega gaccactntt eggatgeana
> 240 cagcaagagg ctttattggg aatnogggta cccgggcgac ncantotato ngaagactgg
> 242 cgttattttt tttntttttt ttttttgaat tnccngggac anccnnctna gnntanctnc
                                                                            300
                                                                            333
> 244 nctntnnct nccctcctta cttctnntnt ntn
```

RAW SEQUENCE LISTING ERROR SUMMARY PATENT APPLICATION: US/10/660,893

DATE: 07/01/2004 TIME: 13:09:24

Input Set : A:\New Link, 0003.ST25.txt Output Set: N:\CRF4\07012004\J660893.raw

ase Note:

of n and/or Xaa have been detected in the Sequence Listing. Please review the uence Listing to ensure that a corresponding explanation is presented in the <220> <223> fields of each sequence which presents at least one n or Xaa.

#:4; N Pos. 3,4,8,350,358,359,360,361,362,363,364,365,366,367,368,369

#:4; N Pos. 370,371,372,373,374 #:5; N Pos. &,11,106,416,168,179,204,211,224,231,254,2/2,275,282,285,286 #:5; N Pos. 205,292,293,296,299,301,304,306,307,308,311,326,327,328,3/31

#:5; N Pos: 333

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/660,893

DATE: 07/01/2004 TIME: 13:09:24

Input Set : A:\New Link, 0003.ST25.txt
Output Set: N:\CRF4\07012004\J660893.raw

7 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:0
7 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:300
9 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:4 after pos.:360
34 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0
36 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:60
38 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:120
40 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:180
42 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:240
44 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:240